

Disseminated Tuberculosis in Interferon γ Gene-disrupted Mice

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Summary

The expression of protective immunity to *Mycobacterium tuberculosis* in mice is mediated by T lymphocytes that secrete cytokines. These molecules then mediate a variety of roles, including the activation of parasitized host macrophages, and the recruitment of other mononuclear phagocytes to the site of the infection in order to initiate granuloma formation. Among these cytokines, interferon γ (IFN- γ) is believed to play a key role in these events. In confirmation of this hypothesis, we show in this study that mice in which the *IFN- γ* gene has been disrupted were unable to contain or control a normally sublethal dose of *M. tuberculosis*, delivered either intravenously or aerogenically. In such mice, a progressive and widespread tissue destruction and necrosis, associated with very high numbers of acid-fast bacilli, was observed. In contrast, despite the lack of protective immunity, some DTH-like reactivity could still be elicited. These data, therefore, indicate that although IFN- γ may not be needed for DTH expression, it plays a pivotal and essential role in protective cellular immunity to tuberculosis infection.

Current murine models of experimental *Mycobacterium tuberculosis* infection indicate that the emergence of acquired immunity to this organism is mediated by populations of both class I and II MHC-restricted T lymphocytes, which secrete cytokines that result in the activation of parasitized macrophages and promotion of the granulomatous response (1–4). The cytokine IFN- γ , which is an effective inducer of antimicrobial mechanisms in several systems (5, 6) has been shown to inhibit the growth of mycobacteria in vitro (7–9), but its role in vivo has yet to be precisely defined. In this regard, we report here that mice in which the *IFN- γ* gene has been disrupted (*IFN- γ* gene knockout mice [GKO mice]), develop a fatal, disseminated form of disease when inoculated with a normally sublethal inoculum of *M. tuberculosis*. These results indicate, therefore, that IFN- γ plays a pivotal role in the expression of protective immunity to this infection in the mouse. Moreover, this new immunodeficient mouse model may prove highly useful in the evaluation of therapeutic intervention strategies in the severely immunocompromised host.

Materials and Methods

Mice. GKO mice were generated as described previously (10). Briefly, a normal *IFN- γ* allele in mouse embryonic stem cells was replaced with a defective gene using a targeted vector which introduced a termination codon after the first 30 amino acids of the

mature IFN- γ protein. The altered stem cells were injected into C57BL/6J blastocysts and transmitted via the germline. Heterozygous offspring of the chimeras were intercrossed to generate mice homozygous for the altered (GKO) and wild type (WT) allele. The GKO mice were previously characterized as normal in terms of spleen and thymus cell number and expression of CD3, B220, CD4, and CD8 surface markers, and were shown to be incapable of IFN- γ secretion (10).

Experimental Infections. The Erdman strain of *M. tuberculosis* was grown in Proskauer Beck medium containing 0.01% Tween 80 to mid-log phase. Mice were infected intravenously via a lateral tail vein with an inoculum of 10^5 *M. tuberculosis* suspended in 0.2 ml PBS. For airborne infections, mice were placed in the exposure chamber of an airborne infection apparatus (Glas-col Inc., Terre Haute, IN). The nebulizer compartment was filled with 10 ml of a suspension of *M. tuberculosis* at a concentration previously calculated to provide an uptake of ~ 50 viable bacilli within the lungs over a 30-min exposure (1, 2).

The numbers of viable bacteria in target organs was followed against time by plating serial dilutions of whole organ homogenates on nutrient Middlebrook 7H11 agar (GIBCO BRL, Gaithersburg, MD) and counting bacterial colony formation after 21 d incubation at 37°C in humidified air.

Histological Analysis. Tissues were fixed in 10% formal saline, set in paraffin blocks, sectioned, and stained using the Ziehl-Neelsen method to visualize acid-fast bacilli. For electron microscopy, tissues were fixed in 1% glutaraldehyde in HBSS, postfixed in 1% OsO₄, dehydrated in acetone, and embedded in Spurr's resin. Sections were cut and stained in 1% uranyl acetate and Reynold's lead.

Results

In this study, we first wished to ensure that T cells in the GKO mice were in a similar physiological state as the WT controls. Flow cytometric staining of T cells for the CD44 marker, which characteristically changes on activated cells (11), was found to be expressed similarly in both groups of mice. In addition, in assurance of the effectiveness of the gene disruption, abundant mRNA message encoding IFN- γ was detected in the tissues of infected WT controls, but was completely absent in GKO mice (data not shown).

Course of Infections. The course of tuberculosis infection in the GKO and WT mice was followed against time after delivery by two separate routes. In the first, animals were infected intravenously with a normally sublethal dose (10^5 bacilli) of the virulent Erdman strain of *M. tuberculosis*. As shown in Fig. 1 a, evidence of control and containment of the infection in the primary target organs (liver, spleen, and lungs) of WT mice was clearly evident from day 10 onwards. In contrast, however, the infection grew progressively in the GKO mice, reaching very high numbers. The experiment was humanely curtailed on day 28, when remaining GKO mice showed visible signs of severe illness.

A notable observation in the GKO mice was the presence of large numbers of bacteria in the kidney and bone marrow, as well as an obvious bacteremia, indicating severe dissemi-

nation of the infection. In contrast, no bacteria were detected in these tissues in the control animals.

In a second experiment, mice were infected aerogenically with a low dose ($\sim 5 \times 10^1$) of bacteria using an aerosol generator (Fig. 1 b). In the WT control animals, containment of the pulmonary infection was evident after 30–40 d, consistent with our previous observations (12). In addition, some degree of hematogenous seeding to the spleen and liver was observed; again, this is consistent with earlier reports (1). In the GKO mice, however, there was no containment of the infection, which grew progressively. When the experiment was curtailed on day 50 of the experiment, bacterial numbers in the lungs had reached over 10^9 , an extraordinary tissue load. Again, bacteria could be harvested from the kidney, bone marrow, and blood of the GKO mice, but not from these tissues in control animals.

Histology of Infection. Previous studies have demonstrated that emergence of the protective immune response to intravenous infection peaks ~ 2 wk after inoculation (1). At this point both WT and GKO mice showed evidence of some initiation of the granulomatous response, with small perivascular accumulations of mononuclear cells, associated with a few acid-fast bacilli, visible in target organs such as the liver. By week 4, however, the GKO mice exhibited spectacular histopathology, with multifocal necrotic areas containing vast numbers of intralésional acid-fast bacteria evident throughout tissue sections of the spleen, liver, lungs, and kidneys. Fig. 2 shows one such area in the liver, in which a very large halo of bacteria surrounded a large area of necrosis and nuclear debris. Beyond this halo, a few neutrophils, eosinophils, and histiocytes could be seen, but any evidence of mononuclear cell accumulation was completely absent. Similar widespread damage was seen in the lungs, in which there was substantial interstitial hemorrhaging, with only a sparse inflammatory cell infiltrate comprised primarily of neutrophils and eosinophils. In addition, there were bacterial clumps evident throughout the alveoli and in the perivascular areas.

Similar pathology was observed by electron microscopy (Fig. 3). In WT control mice infected 4 wk previously by the aerosol route, discrete, well-formed accumulations of mononuclear cells delineated the main sites of infection. Most of the bacteria within the cells of the granulomas were degraded. In contrast, the lungs of GKO mice were almost totally devoid of mononuclear cells. Instead, the tissue had been extensively infiltrated by neutrophils and eosinophils. The lung tissue had lost its cohesive structure, and exhibited advanced caseous necrosis and multiple foci of bacterial growth. A few bacterial clumps were extracellular, after apparent degradation of their host cell, but the majority had clearly been engulfed by eosinophils. These bacteria appeared to be situated within membrane-bound vacuoles, many of which had fused with the lysosomal granule contents of the cell, although with no apparent damage to the bacilli. Recently, *Mycobacterium* has been shown to enter eosinophils (13), and a role for granulocytes in host defense has been proposed (14). The present results clearly indicate, however, that eosinophils alone are not capable of limiting the growth of *M. tuberculosis*.

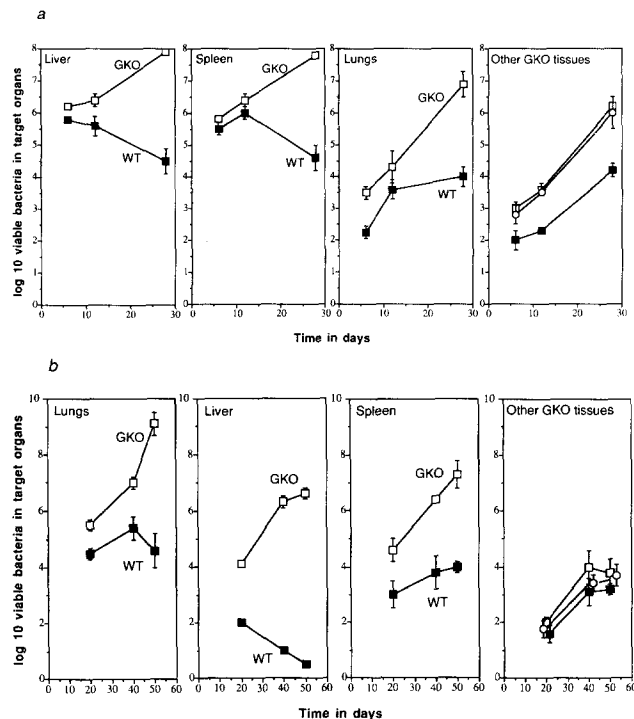


Figure 1. Course of *M. tuberculosis* (strain Erdman) infections in GKO and WT (control) mice. (a) Intravenous inoculation with 10^5 viable bacilli. The course of infection in GKO (\square) and WT control mice (\blacksquare) was followed against time in target organs, and in tissues not normally infected in immunocompetent mice (right: GKO mice; kidney [\square], bone marrow [\circ], and blood [\blacksquare]). Data shown is mean of bacterial counts from four mice, \pm SEM. (b) Course of infection in target organs after the deposition of ~ 50 – 80 viable bacilli in the lungs. Symbols for other infected GKO tissues are as above.

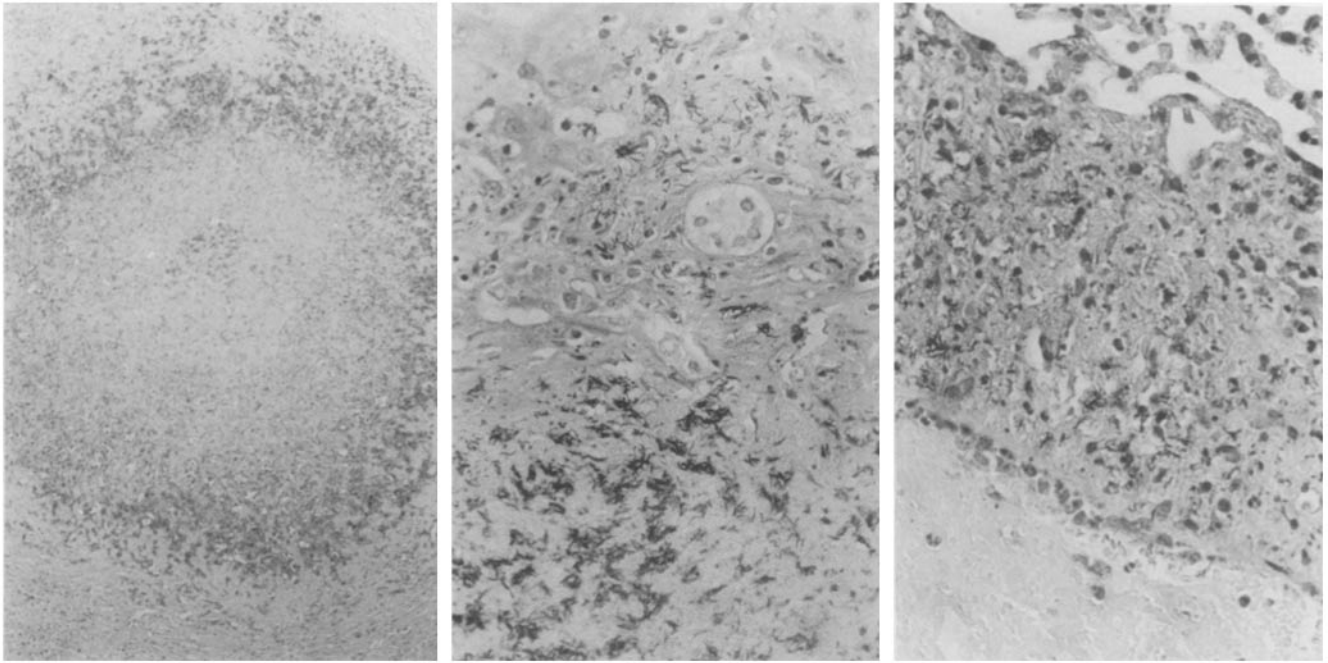


Figure 2. Representative tissue histology obtained from GKO mice 4 wk after infection. Similar patterns of widespread tissue destruction and florid bacterial dissemination were observed after either infection route. (*Top*) A massive halo of bacteria surrounding a large area of complete tissue destruction in the liver; Ziehl-Neelsen stain. $\times 40$. (*Center*) Edge of halo showing presence of a few granulocytes and histiocytes, but complete absence of any mononuclear cell response. $\times 160$. (*Bottom*) Perivascular/parenchymal tissue in lungs containing numerous large clumps of mycobacteria. $\times 160$.

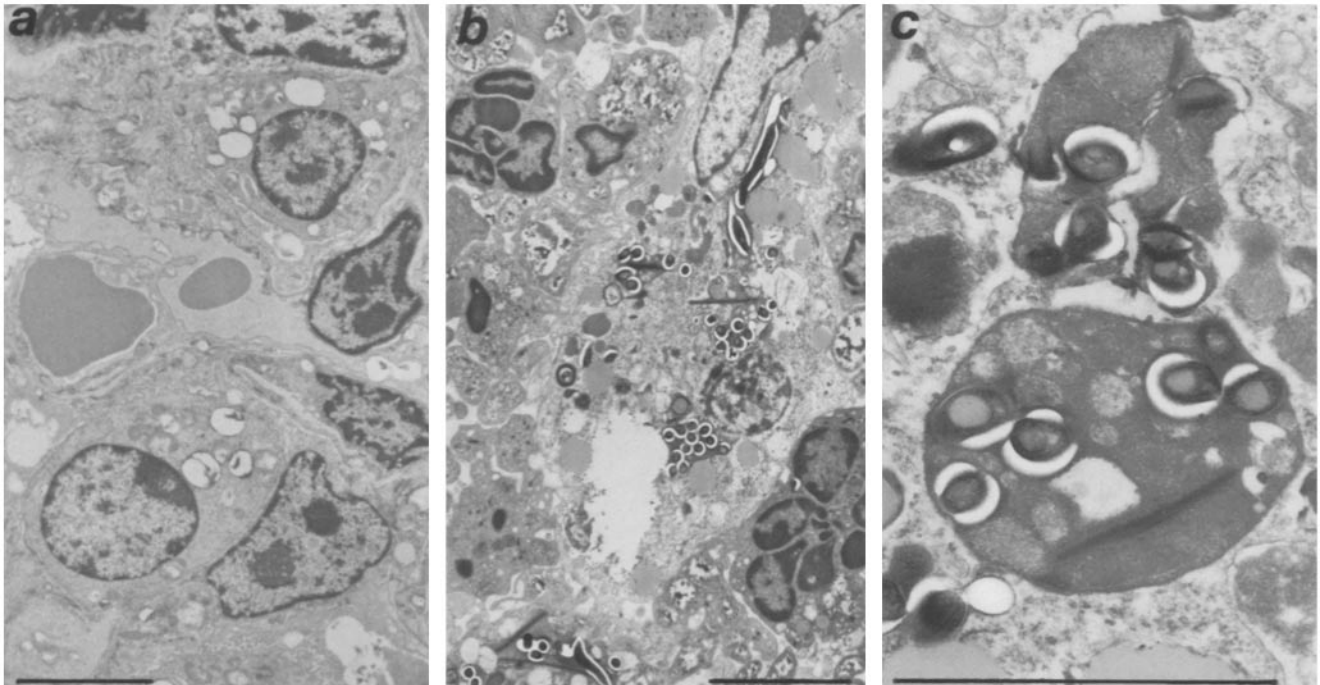


Figure 3. Electron microscopic examination of lung tissues 4 wk after aerogenic infection. (*a*) Discrete granulomatous formation in WT control mice. Some monocytes possess obvious vacuoles, a few of which may contain bacterial remnants. The size bar at the bottom of this photograph corresponds to $5 \mu\text{m}$. (*b*) An area of extensive lung tissue necrosis and bleeding in GKO mice. Scattered granulocytes, and large rafts of extracellular bacteria are evident. Size bar = $5 \mu\text{m}$. (*c*) Phagocytosis of several mycobacteria by an eosinophil in the lung of a GKO mouse. Size bar = $2.5 \mu\text{m}$.

Discussion

This study shows that mice in which the gene for the molecule IFN- γ has been disrupted, a normally sublethal inoculum of *M. tuberculosis*, delivered by either the intravenous or aerogenic routes, grew progressively to lethal levels. Histological examination of these mice revealed widespread caseous necrosis throughout the major target organs such as the lungs and liver, with no evidence of a surviving mononuclear cell response at a time when bacterial loads reached very high levels.

These data convincingly demonstrate, therefore, that mice lacking a functional gene for IFN- γ are totally unable to contain and control a virulent *M. tuberculosis* infection. Since we were able to observe the beginnings of a mononuclear cell granulomatous response occurring early during the course of the infection in the GKO mice, it seems reasonable at this time to hypothesize that the central deficiency in these animals lies in their inability to adequately activate both infected macrophages and arriving monocytes to halt the progressive growth of the infection. These data suggest in turn, therefore, that other cytokines can mediate the initiation of the granuloma, but that infiltrating phagocytes, in the absence of IFN- γ -mediated activation, become heavily infected and destroyed leading to the observed widespread tissue necrosis.

In this regard, the cytokine TNF has also been implicated in having a key role in such mechanisms. Kindler et al. (15) found that neutralizing Ab to TNF disrupted the architecture of granulomas and increased bacterial numbers in mice infected with the avirulent BCG strain of *M. bovis*. Similarly, Amiri et al. (16) observed that TNF could reconstitute the granulomatous response to *Schistosoma* eggs in *scid* mice. In this latter model, however, such mice are known to retain NK cells that can also be a source of IFN- γ . Collectively, therefore, it seems reasonable to hypothesize that, while TNF may clearly contribute in some way to the formation of the granuloma, IFN- γ appears to be essential for the successful retention of its integrity, including the expression of bactericidal mechanisms.

The expression of cutaneous sensitivity to skin test reagents containing mycobacterial antigens (DTH reactions) is also believed to represent a type of granulomatous response in sensitized individuals, involving as it does the local accumulation of monocytes (17). Although this reaction remains a mainstay in the clinical diagnosis of tuberculosis, the basis of the

reaction, including the role of cytokines such as IFN- γ , still lacks clear definition. In this regard, both WT and GKO mice mounted significant swelling responses to skin test antigens after footpad inoculation on day 12 of the intravenous infection (0.43 ± 0.10 mm in WT, 0.67 ± 0.37 mm in GKO). Whereas these reactions followed the kinetics of characteristic 24-h DTH responses, we are unable at this time to provide hard histological evidence that these reactions represented true DTH. Should this subsequently be proven to be the case however, then these data would indicate that whereas the loss of the ability to secrete IFN- γ has a devastating effect on protective immunity against *M. tuberculosis*, it may not be essential to the DTH response in which other cytokines such as TNF, migrating inhibitory factor, and IL-8 (15, 18–20) released by macrophages or T cells may be more important. This in turn provides evidence for a possible dissociation between protection and DTH, although it does not necessarily mean that separate T cell populations are involved.

Evidence for a low state of activation of macrophages harvested from GKO mice infected with *M. bovis* BCG has been reported previously (10). Such cells had low levels of reactive oxygen and nitrogen radical production, and were poor expressers of class II MHC molecules. It is of no surprise, therefore, that such cells were incapable of controlling the proliferation of *M. tuberculosis*, and hence were destroyed. Furthermore, many bacteria were present (especially in lung tissues) in areas of extensive caseous necrosis, a pathology not previously believed (17) to exist in the mouse. This model, and another recently published report using β_2 -microglobulin gene-disrupted mice (3), refutes this earlier contention.

In conclusion, the model described both here and in a parallel publication (21) may provide useful new information in the evaluation of several important areas. In particular, from the perspective of infectious disease, it provides an excellent new model whereby the precise role of IFN- γ in the expression of protective immunity to a variety of microorganisms can be thoroughly investigated. In addition, as a model of the severely immunocompromised host, the GKO mouse may prove very useful for the evaluation and testing of new strategies of immunotherapy and chemotherapy of tuberculosis, and other intracellular infections.

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